SOC-GA 2332 Intro to Stats Lab 6

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10/11/2024

Part 1: Gauss-Markov Assumption and Strict Exogeneity

• In lecture, we talked about Gauss-Markov Assumption and the fact that the OLS estimator is the Best Linear Unbiased Estimator (BLUE) if the assumptions are met. For a data generating process (DGP):

$$Y_i = \beta_0 + \beta_1 X_i + \epsilon_i$$

- Zero conditional mean: in this population, $\mathbb{E}(\epsilon_i|X) = 0$. We also call this the strict exogeneity assumption. This means that, no matter which value X takes, the expectation of ϵ_i associated with this X value will be 0. If the assumption is met, the following statements will be true:
 - $-\mathbb{E}(\epsilon_i) = 0$. This is because of the Law of Iterated Expectations (detailed explanations here): $\mathbb{E}(\epsilon_i) = \mathbb{E}(\mathbb{E}(\epsilon_i|X)) = \mathbb{E}(0) = 0$
 - $-\epsilon_i$ is independent of X. In other words, ϵ_i is not a function of X, otherwise $\mathbb{E}(\epsilon_i|X) = \mathbb{E}(f(X)|X) = f(X) \neq 0$
- The independence between ϵ_i and X, if violated, would produced a biased estimation. That is, if we sample from this population and derive $\hat{\beta}_1$, $\mathbb{E}(\hat{\beta}_1) \neq \beta_1$.
 - This will be part of your assignment 2.
 - You don't even have to sample from the population. You can see this biasedness by creating a population where ϵ_i and X are not independent, and when you regress Y_i on X, your derived $\hat{\beta}_1$ will be very different from β_1 .
 - How to force ϵ_i to be dependent on X? You may create ϵ_i to be any function of X. For example, $\epsilon_i = aX_i + b + N(0,1)$ $(N(\mu, \sigma)$ means a normal distribution with mean μ and SD σ).
 - In reality, this is called omitted variable bias.

Part 2: Important Properties of OLS Estimation

• OLS minimizes the sum of the square of the error term

$$\underset{\hat{\beta}_0, \, \hat{\beta}_1}{\operatorname{argmin}} f(\hat{\beta}_0, \hat{\beta}_1) = \sum_{i=1}^n \left(Y_i - \hat{\beta}_0 - \hat{\beta}_1 X_i \right)^2$$

• We use partial derivative for the solution

$$\begin{split} \frac{\partial f(\hat{\beta_0},\hat{\beta_1})}{\partial \hat{\beta_0}} &= 0 \\ \text{which gives: } \sum_{i=1}^n \left(Y_i - \hat{\beta_0} - \hat{\beta_1} X_i\right) = \sum_{i=1}^n e_i = 0 \\ \frac{\partial f(\hat{\beta_0},\hat{\beta_1})}{\partial \hat{\beta_1}} &= 0 \end{split}$$
 which gives:
$$\sum_{i=1}^n X_i \left(Y_i - \hat{\beta_0} - \hat{\beta_1} X_i\right) = \sum_{i=1}^n X_i e_i = 0 \end{split}$$

- The two facts, $\sum_{i=1}^{n} e_i = 0$ and $\sum_{i=1}^{n} X_i e_i = 0$, are forced to be true in OLS estimation
 - e_i does not have life on its own. It has its meaning and value through $\hat{\beta}_0$ and $\hat{\beta}_1$
 - $\sum_{i=1}^{n} X_i e_i = 0$ forces the covariance between e_i and X_i to be 0. But this does not imply independence.

Part 3: Multivariate Regression & Interaction with One Dummy

Dummies

- For categorical variables, we create dummies or convert them to 0 or 1 dummies when we want to include them in a regression model
- Note that for a categorical variable that have n categories, the regression model will only have n-1 dummies or categorical variable predictors, because the n^{th} dummy is redundant given that if an observation does not belong to any of the n-1 category, then it must be in the n^{th} category
- We call the left-out category the reference category
- Question: what if we include all n categories?
- You should always interpret your model coefficients with the reference category in mind. This could
 get complicated when you have multiple dummy variables, especially when they are interacted in your
 model

In the case of the dummies representing "race" in the earnings_df that we will be using today, we have:

Category	$Dummy_1(black)$	$\overline{Dummy_2(other)}$
White	0	0
Black	1	0
Other	0	1

Exercise (from Lab 5)

- 1. Import earnings_df.csv to your environment. Perform the following data cleaning steps: (1) If age takes the value 9999, recode it as NA; (2) Create a new variable female that equals 1 when sex takes the value female, and equals to 0 otherwise; (3) Create a new variable black that equals 1 when race is black and equals to 0 otherwise; (4) Create a new variable other that equals to 1 when race is 'other' and 0 otherwise.
- 2. Use the describe() function from the psych package to generate a quick descriptive statistics of your data.
- 3. Now, estimate the following models and display your model results in a single table using stargazer(m_1, m_2, ..., m_n, type="text").

```
(1) Model 1: earn \sim age (baseline)
```

- (2) Model 2: earn \sim age + edu
- (3) Model 3: earn \sim age + edu + female
- (4) Model 4: $earn \sim age + edu + female + race$
- (5) Model 5: $earn \sim age + edu + female + race + edu*female$
- 4. Write down your prediction equation for Model 5
- 5. In Model 5, holding other variables constant, what will be the predicted difference in estimated mean earnings for a white man and a white women?
- 6. Holding other variables constant, what will be the predicted difference in estimated mean earnings for a white women and a black women?
- 7. Holding other variables constant, what will be the predicted difference in estimated mean earnings for a white man and a black women?

```
## read data
earnings_df <- read.csv("data/earnings_df.csv", stringsAsFactors = F)</pre>
## recode age
earnings_df <-
  earnings_df %>%
  mutate(age = case_when(
    age > 9000 ~ NA,
    .default = age
  ))
## recode female
earnings_df <- earnings_df %>%
  mutate(female = case_when(
    sex == "female" ~ 1,
    .default = 0))
## base R way of doing it
earnings_df$female <- 0</pre>
earnings_df[earnings_df$sex=="female", "female"] <- 1</pre>
## create black and other
earnings_df <-
  earnings_df %>%
  mutate(black = case when(
    race == "black" ~ 1,
    .default = 0
  )) %>%
  mutate(other = case_when(
    race == "other" ~ 1,
    .default = 0
  ))
m1 \leftarrow lm(earn \sim age,
         data = earnings_df)
m2 \leftarrow lm(earn \sim age + edu,
         data = earnings_df)
m3 <- lm(earn ~ age + edu + female,
```

## ##	=========	=======		========		
::: ## ##	Dependent variable:					
####		(1)	(2)	earn (3)	(4)	(5)
•••	age				0.158***	
	edu				4.477*** (0.112)	
	female				-20.572*** (0.565)	
	black				-2.307*** (0.623)	
•	other				-0.767 (1.137)	
•••	edu:female					-3.128*** (0.199)
####		(1.917)	(1.817)	(1.207)	26.429*** (1.230)	
#	Observations Adjusted R2	980 0.009	980 0.399	980 0.743		980 0.797
	Note:	=======	=======		=======; ; **p<0.05;	***p<0.01

Part 4: Interaction with Two Dummy Variables

Given the following modeling result, please answer the questions.

Table 1:

	$Dependent\ variable:$		
	earn		
college	6.129***		
	(0.187)		
black	-2.773^{***}		
	(0.183)		
college:black	1.496***		
	(0.340)		
Constant	15.077***		
	(0.102)		
Observations	5,000		
\mathbb{R}^2	0.290		
Adjusted R ²	0.289		
Residual Std. Error			
F Statistic	$679.910^{***} (df = 3; 4996)$		
Note:	*p<0.1; **p<0.05; ***p<0.05		

- 1. What will be the predicted difference in estimated mean earnings for a white person with a college degree and a black person with a college degree? Whose earnings will be higher?
- 2. What will be the predicted difference in estimated mean earnings for a white person with a college degree and a black person without a college degree? Whose earnings will be higher?
- 3. How to interpret the interaction coefficient?
- 4. How to interpret the intercept?

Part 5: Visualize Modeling Results

earn 1.0000000 0.624842532 0.100854764

Correlation Matrix

- Sometimes, it is helpful to get an understanding of how variables are linearly related to each other. This is useful in identifying multicollinearity in the data.
- However, since correlation only works with numeric data, you need to remove non-numeric and irrelevant variables before you calculate the correlation matrix using cor().

```
## remove non-numeric variables
earnings_df_cat <- earnings_df %>%
   select(-female, -black, -other, -sex, -race, -unique_id)

## correlation matrix
## set use = "complete.obs" to ignore observations with NAs
cor(earnings_df_cat, use = "complete.obs")

## earn edu age
```

```
## edu 0.6248425 1.000000000 0.002734791
## age 0.1008548 0.002734791 1.000000000
```

stargazer and Multi-category Dummies

##							
## ======= ##	Dependent variable:						
##	-						
##		earn	4-3				
## ##	(1)	(2)	(3)				
;# ;# age	0.158***	0.156***	0.158***				
! #	(0.022)	(0.019)	(0.022)				
# #							
## edu	4.477***						
# #	(0.112)	(0.143)	(0.142)				
##							
## female	-20.572***						
## ##	(0.565)	(1.311)	(0.565)				
+# ## black	-2.307***	-2.385***	-2.023				
##		(0.557)					
# #			,				
## other	-0.767	-0.946	-2.682				
##	(1.137)	(1.017)	(3.000)				
##							
## edu:female		-3.128***					
#		(0.199)					
## ## edu:black			-0.048				
## edu:black ##			(0.243)				
;# !#			(0.240)				
## edu:other			0.336				
##			(0.484)				
##							
## Constant	26.429***	16.974***	26.449***				

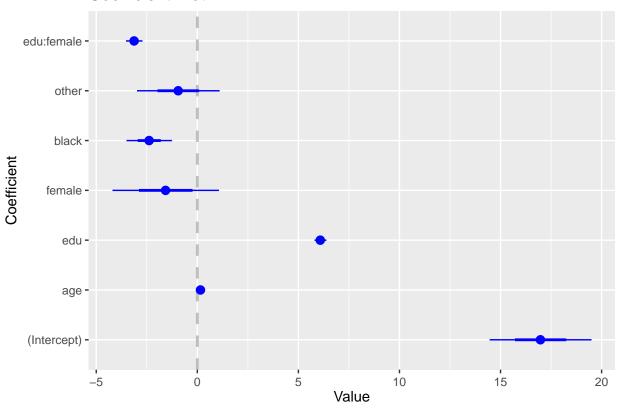
```
(1.230)
                         (1.254)
                                  (1.322)
##
##
##
                 980
                          980
                                   980
## Observations
## Adjusted R2
                0.746
                          0.797
                                  0.745
## ===============
## Note:
                *p<0.05; **p<0.01; ***p<0.001
```

3. Coefficient Plots

- Coefficient plot visualizes the coefficients with it's confidence intervals. You can plot it easily using coefplot() from the coefplot package. There are also other packages that visualize coefficients.
- You can also visualize them on your own by putting the coefficients together in a dataframe.

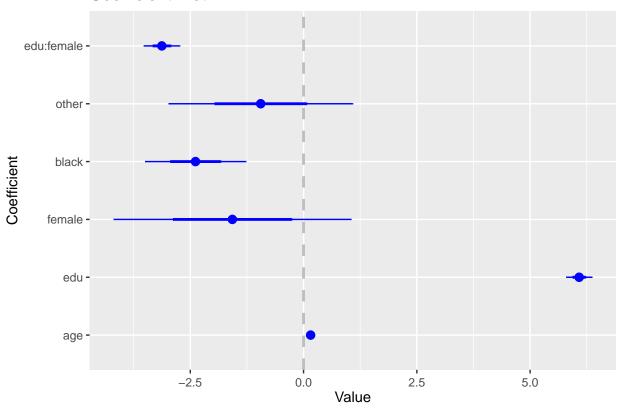
defualt coefficient plot
coefplot(m5)

Coefficient Plot



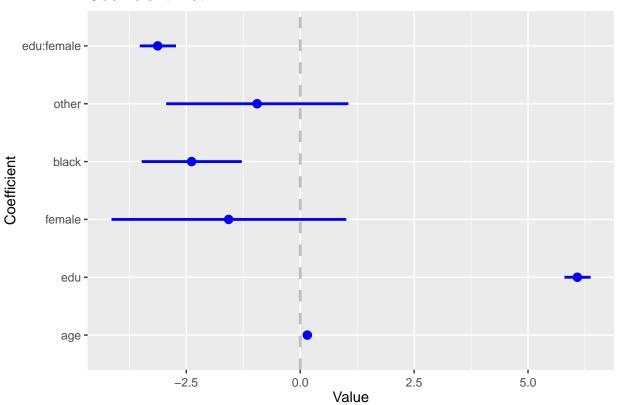
remove the intercept from the plot
coefplot(m5, intercept = F)

Coefficient Plot



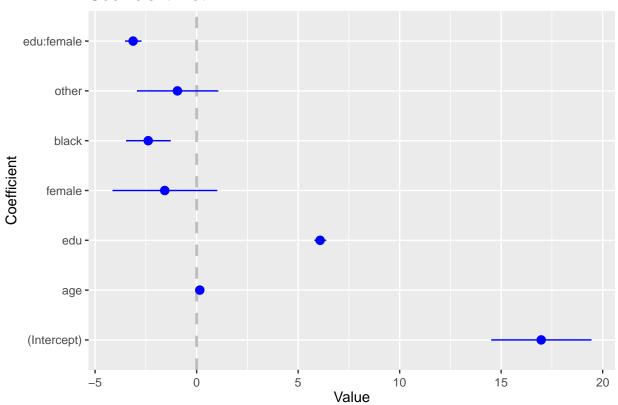
```
## the default innerCI is 1, which is 1 se around the point estimate
## the default outerCI is 2, which is 2 se around the point estimate
## you can set both to 1.96, which is the 95% confidence interval of betas
coefplot(m5, intercept = F, innerCI = 1.96, outerCI = 1.96)
```

Coefficient Plot

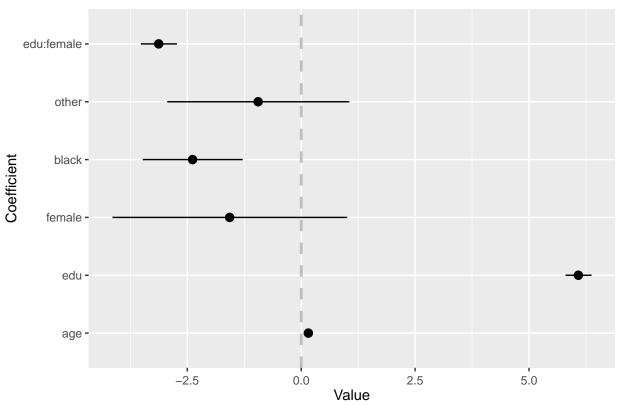


or only keep the outerCI = 1.96
coefplot(m5, intercept = T, innerCI = F, outerCI = 1.96)

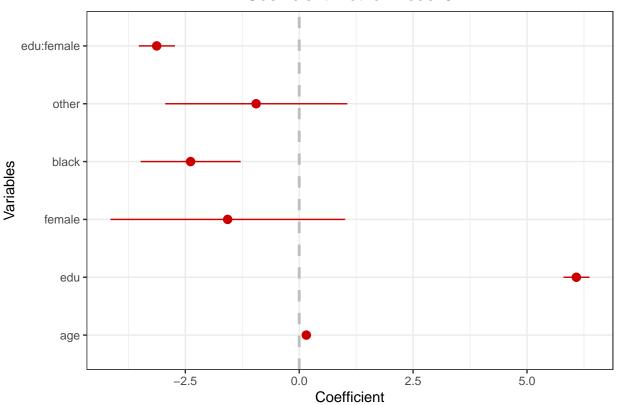
Coefficient Plot



Coefficient Plot for Model 5



Coefficient Plot for Model 5



4. Plot Predicted Effects

- We can visualize the predicted effects of key predictors using the predict() function in base R.
- The idea behind this task is to first create a dataframe with values of all the predictors included in the model, with only the value of your predictor(s) of interest vary within the possible range, whereas other predictors held at their mean.
- For example, if we want to examine the effect of **education and gender** on earnings, we create a dataframe with a variable **edu** that varies from 0 to 15 with an interval of 1 (so **edu** = 0, 1, 2, ..., 14, 15), because the possible value of **edu** in our data is integers from 0 to 15 (you can use **summary(your_df)** to check value ranges).
- We repeat this number sequence for another time so that we have **each level of education for both** male and female. So we need to generate edu = 0, 1, 2, ..., 14, 15, 0, 1, 2, ..., 14, 15. We use rep(0:15, 2) to generate this number sequence.
- rep(x, times) replicate x (a vector or list) for user-defined times (in our case, times = 2). You can run this in your R console to see what number sequence is returned.
- Then, we generate a dummy variable female that equals to 0 for male and 1 for female.
- To create a dataframe that have the combination of each level of edu and each gender category, we let female = 0 for 16 times, and female = 1 for 16 times, using c(rep(0, 16), rep(1, 16)). You can run this in your R console to see what number sequence is returned.
- For the rest of the predictors, we fix them at their mean. We add na.rm = T in the mean() function to specify how we want to deal with NA values. If you don't include na.rm = T, mean() will return NA if your variable contains NAs.

```
## first, we create a dataframe with all predictor variables
## only the key predictor varies, while the others remain at the mean
pred_IV <- data.frame(edu = rep(0:15, 2)) %>%
                                                       ## first, create a df with values of your key pre
  mutate(female = c(rep(0, 16), rep(1, 16)),
                                                       ## b/c we are looking at interaction effects,
                                                        ## give gender two values, otherwise fix it at me
         age = mean(earnings_df$age, na.rm = T),
                                                     ## fix other variables at mean
         black = mean(earnings_df$black),
         other = mean(earnings_df$other))
rep(0:15,2)
## [1] 0 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 0 1 2 3 4 5
## [26] 9 10 11 12 13 14 15
## examine the df
head(pred_IV, 5)
##
     edu female
                     age black other
## 1
              0 43.26429 0.306 0.068
## 2
              0 43.26429 0.306 0.068
       1
## 3
              0 43.26429 0.306 0.068
## 4
       3
              0 43.26429 0.306 0.068
## 5
              0 43.26429 0.306 0.068
  • Now that we have the dataframe pred_IV ready for predicting the dependent variable (earning), we
    can use the R function predict() to calculate fitted earning using the regression model and the values
    specified in each row in pred_IV. Then, use cbind() to combine this fitted Y value vector with your
    pred_IV for plotting.
## use `predict` to predict the Y
predicted_earning <- predict(m5,</pre>
                                                        ## the model you are using
                                                       ## the df you use for predicting
                              interval = "confidence", ## set CI
                              level = 0.95)
## bind the columns
pred_result <- cbind(pred_IV, predicted_earning)</pre>
## check df
head(pred_result, 5)
##
     edu female
                     age black other
                                           fit
                                                     lwr
                                                              upr
## 1
              0 43.26429 0.306 0.068 22.93127 21.12317 24.73936
## 2
              0 43.26429 0.306 0.068 29.01392 27.46140 30.56644
       1
## 3
              0 43.26429 0.306 0.068 35.09657 33.78940 36.40374
## 4
              0 43.26429 0.306 0.068 41.17922 40.10017 42.25827
       3
## 5
              0 43.26429 0.306 0.068 47.26188 46.38025 48.14350
## plot
pred_result %>%
  mutate(gender = ifelse(female == 0, "Male", "Female")) %>%
                                                                     ## convert dummy to character variab
  ggplot(aes(x = edu, y = fit, group = gender)) +
  geom_line(aes(linetype = gender)) +
                                                                     ## group linetype by gender
                                                                            # add 95% CI
  geom_ribbon(aes(ymin = lwr, ymax = upr, fill = gender), alpha = 0.3) +
  theme_bw() +
  labs(x = "Years of Education",
       y = "Predicted Earnings") +
```

Predicted Earnings by Education and Gender

(Modeled with interaction between education and gender)

